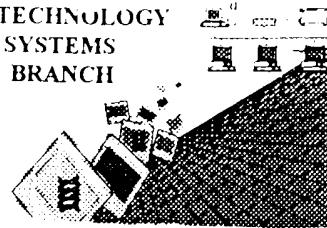


BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: IL USC, 216A
Source: 31PE
Date Processed by STIC: 6/28/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



Does Not Comply
Corrected Diskette Needed

O I P E

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/050,216A

DATE: 06/28/2002
TIME: 11:16:21

Input Set : A:\CORRECTEDSEQUENCE LISTING.txt
Output Set: N:\CRF3\06282002\J050216A.raw

4 <110> APPLICANT: Curtis, Rory A.J., Lora, Jose M.
5 <120> TITLE OF INVENTION: 46798. A Human Matrix Metalloprotease and
6 Uses Therefore
7
8 <130> FILE REFERENCE: MPI2001-014P1RNM
9
10 <140> CURRENT APPLICATION NUMBER: 10/050,216A
C--> 11 <141> CURRENT FILING DATE: 2002-01-16
12
13 <150> PRIOR APPLICATION NUMBER: 60/262,252
14
15 <151> PRIOR FILING DATE: 2001-01-16
16
17 <160> NUMBER OF SEQ ID NOS: 10
18
19 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

746 <210> SEQ ID NO: 10
 747 <211> LENGTH: 1563! found 1563!
 748 <212> TYPE: DNA
 749 <213> ORGANISM: Homo sapiens
 750 <214> : 3
 751 <220> FEATURE:
 752 <221> NAME/KEY: CDS
 753 <222> LOCATION: (1)...(1563)
 755 <400> SEQUENCE: 10
 756 atg gtc gcg cgc gtc ggc ctc ctg ctg cgc gcc ctg cag ctg cta ctg 48
 757 Met Val Ala Arg Val Gly Leu Leu Leu Arg Ala Leu Gln Leu Leu Leu
 758 1 5 10 15
 760 tgg ggc cac ctg gac gcc cag ccc gcg gag cgc gga ggc cag gag ctg 96
 761 Trp Gly His Leu Asp Ala Gln Pro Ala Glu Arg Gly Gly Gln Glu Leu
 762 20 25 30
 764 cgc aag gag gcg gag gca ttc cta gag aag tac gga tac ctc aat gaa 144
 765 Arg Lys Glu Ala Glu Ala Phe Leu Glu Lys Tyr Gly Tyr Leu Asn Glu
 766 35 40 45
 768 caq gtc ccc aaa gct ccc acc tcc act cga ttc agc gat gcc atc aga 192
 769 Gln Val Pro Lys Ala Pro Thr Ser Thr Arg Phe Ser Asp Ala Ile Arg
 770 50 55 60
 772 gcg ttt cag tgg gtg tcc cag cta cct gtc agc ggc gtc ttg gac cgc 240
 773 Ala Phe Gln Trp Val Ser Gln Leu Pro Val Ser Gly Val Leu Asp Arg
 774 65 70 75 80
 776 gcc acc ctg cgc cag atg act cgt ccc cgc tgc ggg gtt aca gat acc 288
 777 Ala Thr Leu Arg Gln Met Thr Arg Pro Arg Cys Gly Val Thr Asp Thr
 778 85 90 95
 780 aac agt tat gcg gcc tgg gct gag agg atc agt gac ttg ttt gct aga 336
 781 Asn Ser Tyr Ala Ala Trp Ala Glu Arg Ile Ser Asp Leu Phe Ala Arg
 782 100 105 110

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/050,216A

DATE: 06/28/2002
TIME: 11:16:21

Input Set : A:\CORRECTEDSEQUENCE LISTING.txt
Output Set: N:\CRF3\06282002\J050216A.raw

784	cac	cg	acc	aaa	atg	agg	cgt	aag	aaa	cgc	ttt	gca	aag	caa	gg	t	aac	384
785	His	Arg	Thr	Lys	Met	Arg	Arg	Lys	Lys	Arg	Phe	Ala	Lys	Gln	Gly	Asn		
786	115				120					125								
788	aaa	tgg	tac	aag	cag	cac	ctc	tcc	tac	cgc	ctg	gtg	aac	tgg	cct	gag		432
789	Lys	Trp	Tyr	Lys	Gln	His	Leu	Ser	Tyr	Arg	Leu	Val	Asn	Trp	Pro	Glu		
790	130				135					140								
792	cat	ctg	ccg	gag	ccg	gca	gtt	cg	gg	gcc	gt	cgc	gcc	gcc	tcc	cag		480
793	His	Leu	Pro	Glu	Pro	Ala	Val	Arg	Gly	Ala	Val	Arg	Ala	Ala	Phe	Gln		
794	145				150					155							160	
796	ttg	tgg	agc	aac	gtc	tca	g	ctg	gag	tcc	tgg	gag	gcc	cca	gcc	aca		528
797	Leu	Trp	Ser	Asn	Val	Ser	Ala	Leu	Glu	Phe	Trp	Glu	Ala	Pro	Ala	Thr		
798					165					170							175	
800	ggc	ccc	gct	gac	atc	cg	ctc	acc	tcc	tcc	caa	ggg	gac	cac	aa	qat		576
801	Gly	Pro	Ala	Asp	Ile	Arg	Leu	Thr	Phe	Phe	Gln	Gly	Asp	His	Asn	Asp		
802					180					185							190	
804	ggg	ctg	ggc	aat	gcc	ttt	gat	ggc	cca	ggg	ggc	gcc	ctg	g	cc	cac	ggc	624
805	Gly	Leu	Gly	Asn	Ala	Phe	Asp	Gly	Pro	Gly	Gly	Ala	Leu	Ala	His	Ala		
806					195					200							205	
808	tcc	ctg	ccc	cg	cg	gaa	g	cg	ca	g	tcc	gac	caa	gat	gag	cg	tgg	672
809	Phe	Leu	Pro	Arg	Arg	Gly	Glu	Ala	His	Phe	Asp	Gln	Asp	Glu	Arg	Trp		
810					210					215							220	
812	tcc	ctg	agc	cg	cg	ggg	cg	cc	aa	ctg	tcc	gt	gt	ctg	g	cc	cac	720
813	Ser	Leu	Ser	Arg	Arg	Gly	Arg	Asn	Leu	Phe	Val	Val	Leu	Ala	His			
814					225					230							235	240
816	gag	atc	ggt	cac	ac	ctt	ggc	ctc	acc	cac	tcg	ccc	g	cc	cg	cg		768
817	Glu	Ile	Gly	His	Thr	Leu	Gly	Leu	Thr	His	Ser	Pro	Ala	Pro	Arg	Ala		
818					245					250							255	
820	ctc	atg	g	cc	tac	tac	aag	agg	ctg	ggc	cg	gac	g	cc	ctg	ctc	agc	816
821	Leu	Met	Ala	Pro	Tyr	Tyr	Lys	Arg	Leu	Gly	Arg	Asp	Ala	Leu	Leu	Ser		
822					260					265							270	
824	tgg	gac	gac	gt	gt	ggc	gt	ca	g	ac	ctg	tat	ggg	aag	cc	cta	ggg	864
825	Trp	Asp	Asp	Val	Leu	Ala	Val	Gln	Ser	Leu	Tyr	Gly	Lys	Pro	Leu	Gly		
826					275					280							285	
828	ggc	tca	gt	ggc	gt	cc	ca	gg	aag	ctg	tcc	act	gac	ttt	gag			912
829	Gly	Ser	Val	Ala	Val	Gln	Leu	Pro	Gly	Lys	Leu	Phe	Thr	Asp	Phe	Glu		
830					290					295							300	
832	acc	tgg	gac	tcc	tac	agc	ccc	caa	gg	agg	cg	cct	gaa	ac	cg	gg		960
833	Thr	Trp	Asp	Ser	Tyr	Ser	Pro	Gln	Gly	Arg	Arg	Pro	Glu	Thr	Gln	Gly		
834					305					310							315	320
836	cct	aaa	ta	tgc	ca	t	tcc	tcc	tac	gat	g	cc	at	ct	gt	ac	gg	1008
837	Pro	Lys	Tyr	Cys	His	Ser	Ser	Phe	Asp	Ala	Ile	Thr	Val	Asp	Arg	Gln		
838					325					330							335	
840	cag	caa	ctg	ta	att	ttt	aaa	ggg	agc	ca	t	tc	tg	gag	gt	gca	g	1056
841	Gln	Gln	Leu	Tyr	Ile	Phe	Lys	Gly	Ser	His	Phe	Trp	Glu	Val	Ala	Ala		
842					340					345							350	
844	gat	ggc	aa	gtc	tca	ga	cc	cgt	cca	ctg	ca	gaa	aga	tgg	gtc	gg		1104
845	Asp	Gly	Asn	Val	Ser	Glu	Pro	Arg	Pro	Leu	Gln	Glu	Arg	Trp	Val	Gly		
846					355					360							365	
848	ctg	cc	cc	cc	aa	cc	tt	gg	at	g	ct	gca	ttg	aa	at	gt	gg	1152

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/050,216A

DATE: 06/28/2002
TIME: 11:16:21

Input Set : A:\CORRECTEDSEQUENCE LISTING.txt
Output Set: N:\CRF3\06282002\J050216A.raw

849	Leu	Pro	Pro	Asn	Ile	Glu	Ala	Ala	Ala	Val	Ser	Leu	Asn	Asp	Gly	Asp	
850	370				375					380							
852	ttc	tac	ttc	ttc	aaa	ggg	ggt	cga	tgc	tgg	agg	ttc	cgg	ggc	ccc	aag	1200
853	Phe	Tyr	Phe	Phe	Lys	Gly	Gly	Arg	Cys	Trp	Arg	Phe	Arg	Gly	Pro	Lys	
854	385				390				395							400	
856	cca	gtg	tgg	ggt	ctc	cca	cag	ctg	tgc	cgg	gca	ggg	ggc	ctg	ccc	cgc	1248
857	Pro	Val	Trp	Gly	Leu	Pro	Gln	Leu	Cys	Arg	Ala	Gly	Gly	Leu	Pro	Arg	
858						405			410							415	
860	cat	cct	gac	gcc	gcc	ctc	ttc	ttc	cct	cct	ctg	cgc	cgc	ctc	atc	ctc	1296
861	His	Pro	Asp	Ala	Ala	Leu	Phe	Phe	Pro	Pro	Leu	Arg	Arg	Leu	Ile	Leu	
862						420			425							430	
864	ttc	aag	ggt	gcc	cgc	tac	tac	gtg	ctg	gcc	cga	ggg	gga	ctg	caa	gtg	1344
865	Phe	Lys	Gly	Ala	Arg	Tyr	Tyr	Val	Leu	Ala	Arg	Gly	Gly	Leu	Gln	Val	
866						435			440							445	
868	gag	ccc	tac	tac	ccc	cga	agt	ctg	cag	gac	tgg	gga	ggc	atc	cct	gag	1392
869	Glu	Pro	Tyr	Tyr	Pro	Arg	Ser	Leu	Gln	Asp	Trp	Gly	Gly	Ile	Pro	Glu	
870						450			455							460	
872	gag	gtc	agc	ggc	gcc	ctg	ccg	agg	ccc	gat	ggc	tcc	atc	atc	ttc	ttc	1440
873	Glu	Val	Ser	Gly	Ala	Leu	Pro	Arg	Pro	Asp	Gly	Ser	Ile	Ile	Phe	Phe	
874						465			470							480	
876	cga	gat	gac	cgc	tac	tgg	cgc	ctc	gac	cag	gcc	aaa	ctg	cag	gca	acc	1488
877	Arg	Asp	Asp	Arg	Tyr	Trp	Arg	Leu	Asp	Gln	Ala	Lys	Leu	Gln	Ala	Thr	
878						485			490							495	
880	acc	tcg	ggc	cgc	tgg	gcc	acc	gag	ctg	ccc	tgg	atg	ggc	tgc	tgg	cat	1536
881	Thr	Ser	Gly	Arg	Trp	Ala	Thr	Glu	Leu	Pro	Trp	Met	Gly	Cys	Trp	His	
882						500			505							510	
884	gcc	aac	tcg	ggg	agc	gcc	ctg	ttc	tga								1563
885	Ala	Asn	Ser	Gly	Ser	Ala	Leu	Phe	*								
886						515			520								

E--> 888

(- 7 -) *- delete* *delete*

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/050,216A

DATE: 06/28/2002
TIME: 11:16:22

Input Set : A:\CORRECTEDSEQUENCE LISTING.txt
Output Set: N:\CRF3\06282002\J050216A.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:489 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:490 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:491 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7
L:493 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:494 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:495 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7
L:498 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:499 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:500 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7
L:503 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:504 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:505 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7
L:508 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7
L:509 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:888 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:888 M:252 E: No. of Seq. differs, <211> LENGTH:Input:1560 Found:1563 SEQ:10